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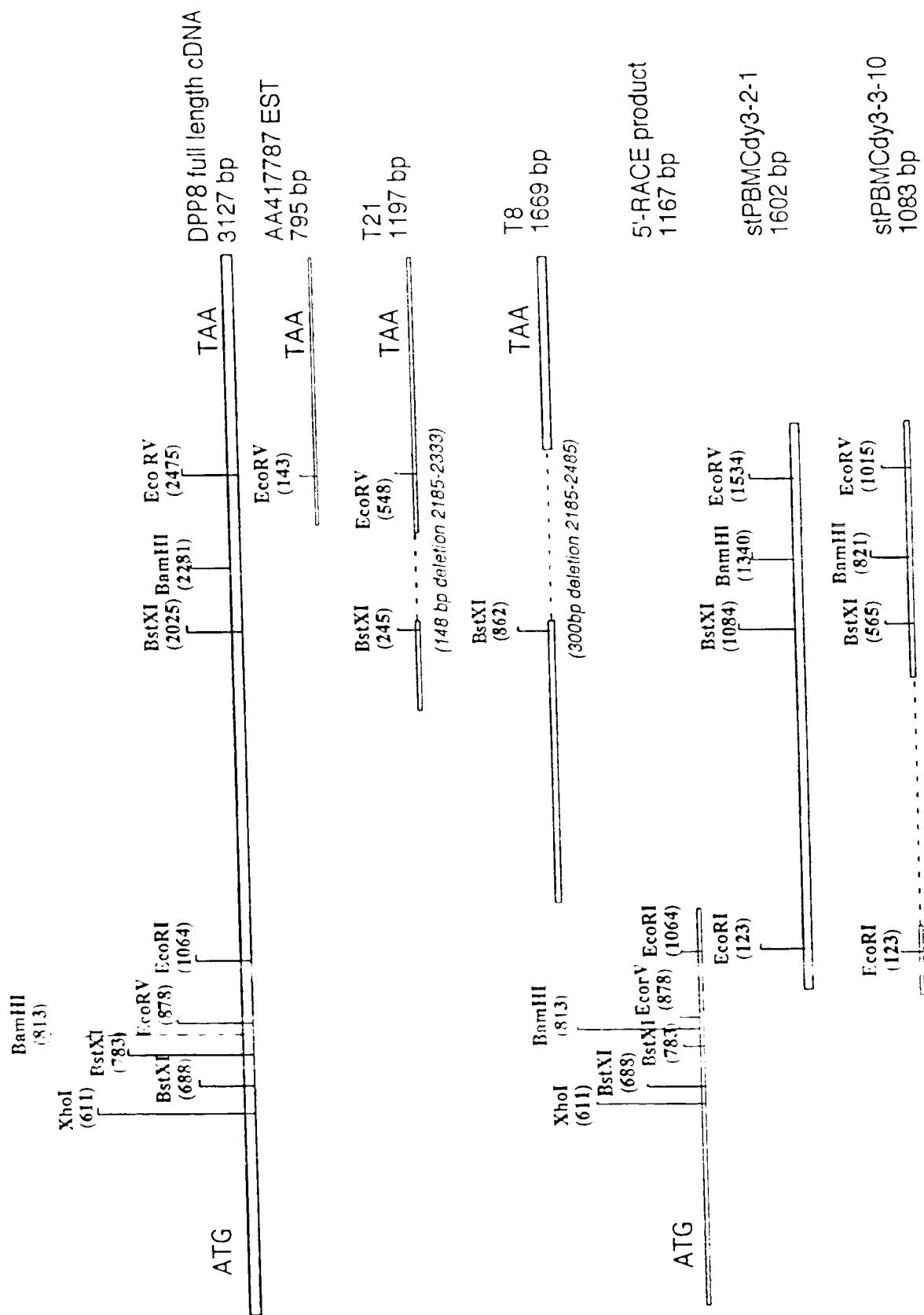


FIGURE 1

1 AGTGCTAAGCTCTGGAGGCCAAGGGCGCTGCTACTGCGCGCGCTGCTTCTTAGTGCGCGGTTGCGCGCTGGTTGTCACCCGGCGCCGGCGAGGA 100
 101 AGCCACTGCAACCAAGGACCGGACTGAGCGGCGGCGACGCTGCAAGCGCGCAAGCGCGCTGCTCATAGCGACGCTGGAGCGTCGGGGCGGGGGGGGA 200
 201 AGGAAATGCAACATGGCAGCAGCAAGAACAGAACAGCTGGCTGTTGAGATAATTGAAACTGCGGACTGTGAGGAGAATATTGAATCACAGGATCGG 300
 KAAAKETEOLGVIEFETADCEENIESQDR 29
 301 CCTAAATGGAGCCCTTTTATGTTGAGCGGTATTCCTGGCTGAGCTTAAAGCTGCTTGGCGATACCGAGAAATATCATGGCTACATGATGGCTAAGG 400
 30 P K L E P F Y V E R Y S W S O L K K L A D T R K Y H G Y H M A K A 63
 401 CACACATGATTCTAGTTGAGAGGAATGATCCAGATGGACCTCATCACAGAACAGATCTATTACCTGGCTAGTCTGGTGAGAACAGAGAAATAC 500
 63 P H D F H F V K R N D P D G P H S D R I Y Y L A M S G E N R E N T 96
 501 ACTCTTTATTCGAAATTCCAAAATCTCAATAGAGCAGCAGCTTAATGCTCTTGGCTGAGCTCTTGGATCTTCAAGGAAACACTGGACTAT 600
 96 L F Y S E I P K T I N R A A V L M L S H K P L L D L F Q A T L D Y 129
 601 CGAAATGATTCGAGAGAAAGAAACTATTAGAGAAAGAAAACCGATTGGAAAGCTGGAAATTGCTCTTACGATTATCAGAACAGGAGTGGACCATTC 700
 130 G M Y S R E E E L L R E R K R I G T V G I A S Y D Y H Q G S G T F L 163
 701 TGTTCAGGGCTAGTGGAAATTATCACGTAAGAACATGGAGGGCCACAAAGGATTAGCGAACACCTTTAAGGCCAACTCTAGTGGAAACTAGTTGTC 800
 163 F Q A G S G I Y H V K D G G G P Q G F T Q Q P R N L V E T S C P 196
 801 CAACATACGGATGGATCCAAAATATGCCCGGCTGATCAGACTGCTGCTTACATAGCGAACAGATTTGGATCTAACATGTAACAGAAA 900
 196 N I R D P K L C P A D P D W I A F I H S N D I V I S N I V T R E 229
 901 GAAGGAGACTCACTTATGACAAATGGCTAGCCAACTGGAGAGATGCCAGATCACGCTGGAGTGGCTACCTTGTCTCAAGAACATTGATA 1000
 230 E R R L T Y V H N E L A N H E D A R S A G V A T F V L Q E E F D R 263
 1001 GATATTCTGGCTATGGTGGTGTCAAAGCTGAAACAACTCCAGTGGTGTAAATTCTAGAATTCTATGAAAGAAATGATGAACTGAGGTGGA 1100
 263 Y S G Y W H C P K A E T T P S G G K I L R I L Y E E N D E S E V E 296
 1101 KATTATTCATTTACATCCCTATGTTGAAACAAAGGAGGGCAGATTCTACCTGGCTTATCTAAACAGGTACAGCAATTCTAAAGTCACCTTAAAGATC 1200
 296 I I H V T S P K L E T R R A D S F R Y P K T G T A N P K V T F K H 329
 1201 TCAAGAAATATGATTGATGCTGAGGAAGGATCATAGATGTCATAGATANGGAACTAATCTAACCTTTGAGATCTTATTTGAGGAGTTGAAATATTTG 1300
 330 S E I M Y D A E G R I I D V I D K E L T I Q P F E I L F E G V E Y I A 363
 1301 CCAGACCTGGATGACTCTGGAGGAAAATATGCTTGGCTACATGACTAGATGGCTCCAGACTGGCTACAGATAGTGTGATCTCACCTGAATTATT 1400
 363 R A G T H P E G K Y A W S T I L L D R S O T R L Q I V L S I P E L F 396
 1401 TATCCGAGTAAAGATGATGTTATGGAAAGGAGACTATTGAGCTAGTCTGGCTGATCTGACCGGCTAAATTATCTATGAGAACACAGACATC 1500
 396 I P V E D D V H E R Q R L I E S V P D S V T P L F I Y E E T T D I 429
 1501 TGGATAAATATCCATGACATCTTCATGTTTCCCAAAGTCAGGAGGAAATTGAGTTTATTTTCCCTCTGAAATGCAAACAGTTGGCTTATT 1600
 430 W I N I H D I F H V F P O S H E E E I E F I F A S E C K T G F R H L 463
 1601 TATACAAAATTACATCTATTAAAGGAAAGCAAATAAAACGTCAGTGGTGGCTGCTGCTCAAGTGTGATTCTGACCTATCTATGAGAACACAGACATC 1700
 463 Y K I T S I L K E S K Y K R S S G G L P A P S D F K C P I K E E I 496
 1701 AGCAATTACAGTGGTGAATGGGAAGTCTGGCCGGCATGGCTAAATCTCAAGTGTGATGAGCTAGAACAGGCTGGTATATTGAGGACCAAAAGAC 1800
 496 A I T S G E W E V L G R H G S N I Q V D E V R R L V Y F E G T K D 529
 1801 TCCCTTCTAGAGCATCACCTGAGCTGAGTTCAGTAAATCTGGAGAGGTGACAGGCTGACTGACCGTGGCTACTCACATTCTGGCTACAGTC 1900
 530 S P L E H H L Y V V S Y V N P G E V T R L T D R G Y S H S C C I S Q 563
 1901 AGCACTGTCACCTCTTAAAGTAAGTATGAACTCACAGAACAGATCCACACTGCTGCTCTTACAGCTATCACGCTGAAAGATGACCAACTTGCA 2000
 563 H C D F F I S K Y S N Q K N P H C V S L Y K L S S P D D P T C K 596
 2001 AACAAAGGAATTGGGCCACCATTTGGATTGAGCTGGAGGCTCTCTGACTATACCTCTCCAGAATTCTCTGAGACTACTGATGATTTACA 2100
 596 T K E F W A T I L D S A G P L P D Y T P P E I S F S E S T T G F T 629
 2101 TCTATGGAGCTCTACAGCCTCATGATCTACACCTGAGAACAAATACCTCTGCTGCTTACATATATGGTGGCTCTGGCTGAGCTGGTGTGAAATA 2200
 630 L Y G H L Y K P H D L Q P G K K Y P T V L F I Y G G P Q V O L V N N 663
 2201 ATGGTTAAAGGACTCAAGTATTCGGCTGAAATACCCCTACGGCTCTCTAGGTATCTGCTGAGTGTAGACACACAGGGATCTCTCACCGAGGGCT 2300
 663 R F K G V K Y F R L N T L A S L G Y V V V V I D N R G S C H R G L 696
 2301 TAAATTGAGCGCTTAAATATAAAATGGCTAAATAGAAATTGAGCATACGGTGAAGGACTCTAACATCTAGCTTCTGATATGATCTTGAC 2400
 696 F F E G A P F Y E R G Q Z E I E D D C V E G C L U Y T L S F T D Y I D 729
 2401 TTAGATGCTGGCATCACCGCTCTGGCTATGGAGGAACTCTGGCTGAGCTTAATGGCTGAGGAGCTGAAATCTAGCTTCTGATATGATCTTGAG 2500
 730 I D R V G T H 3127 G Y L S L H A M L O R S D I T R V A I A G A 763
 2501 CCCACTGACTCTGGATCTCTGATACAGGATACACGGAGCTGTTATGGCTCTCTGGCTGAGGAGCTGAAATGGCTGAGGAGCTGAACTTACAGGATCTGGC 2600
 763 P V T L W I F Y D T G Y T E R Y K H G P D O N E O G Y Y L G S V A 796
 2601 CATGCAAGGAGAAAGTCTCCCTGAAACCAAATGTTACTGCTCTACATGGTTCTGGATGAGAATGCTGAGGAGCTGAACTTACAGGATCTGGC 2700
 796 M Q A E K F P S E P H R L L L N G F L B E N V H F A N T S I L L 829
 2701 AGTTTTTACTGAGGGCTGAGAACCCATATGATCTACAGATCTACGGAGAGACAGCATAAGAGTCTGAACTGGAGAACATTATGAACTG 2800
 830 S F L V R A G K P Y D L Q I Y P O E R H S I R V P E S G E H Y E L K 863
 2801 ATTTTCTGAGGAGCTTCAAGAACCTGGATCACGTTCTGCTCTAAAGTGATATAATTGAGCTGAGGAGCTGAACTCTCTGAGTACACTGGCTA 2900
 863 L L H Y L Q E N L G S R I A A L K V I 882
 2901 TTAAACCAAATGAGGAGGTTAAATCAACAGAAACACAGAAATGATCATCACATTGATACCTGAGAACATCTACTCTGAAATAATGAGG 3000
 3001 CCATCACCGGCTACGGTGTGAGCTGAGCTAATCTAACCTAACCCACAGCTGAAATCAAAATGATACATATTCTGAGAGACCCAGCAATACCCATA 3100
 3101 AGAATTACTA 3127

FIGURE 2

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FIGURE 3
SUBSTITUTE SHEET (RULE 26) RO/AU

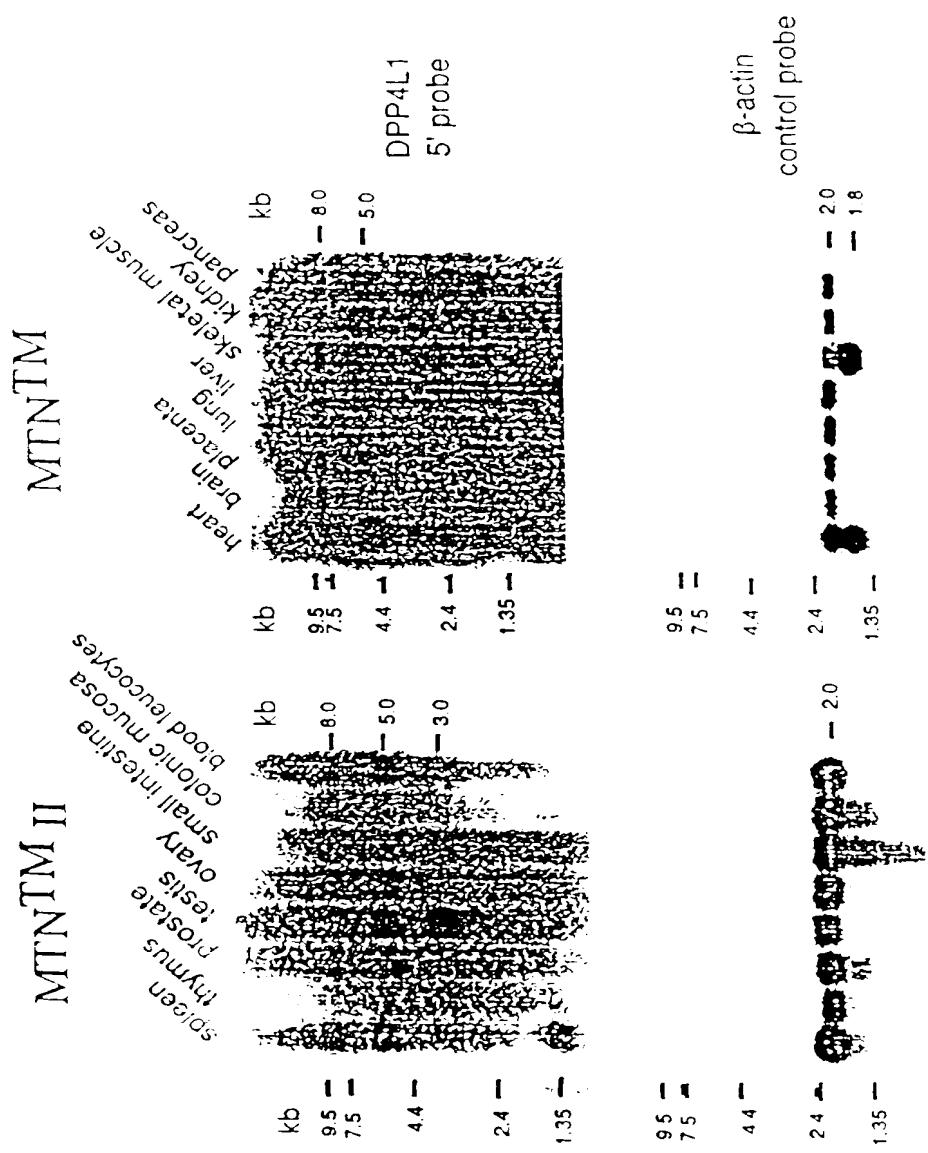


FIGURE 4

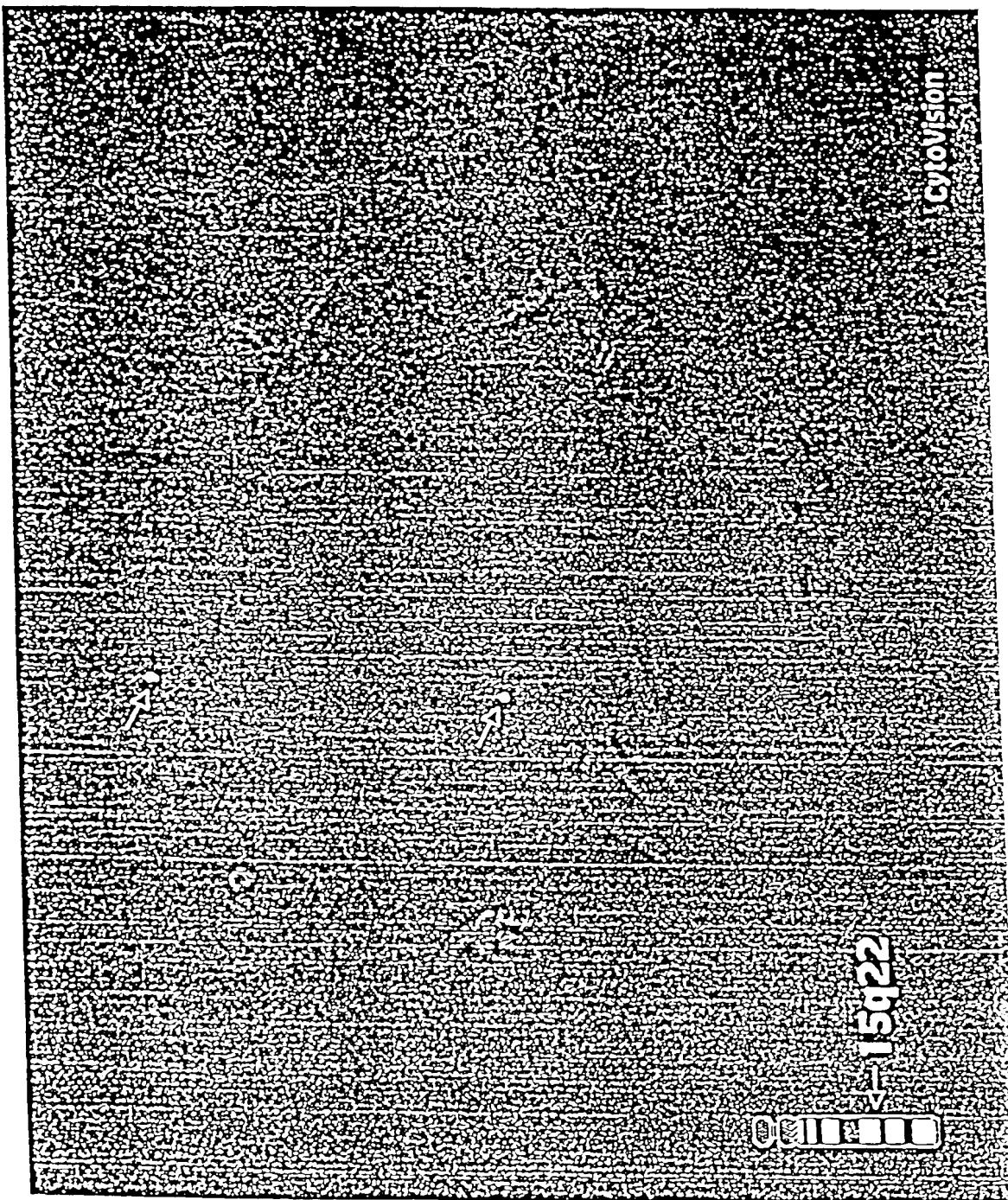


FIGURE 5

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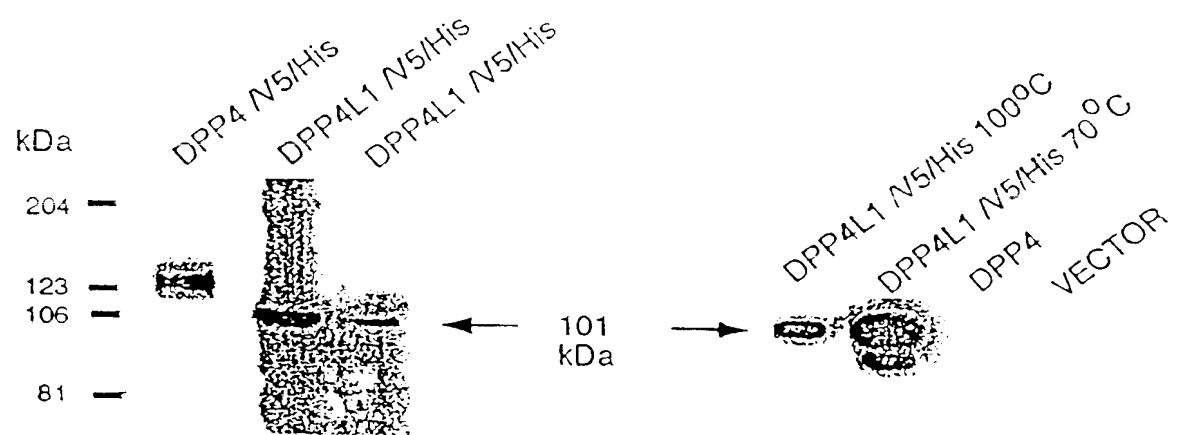


FIGURE 6

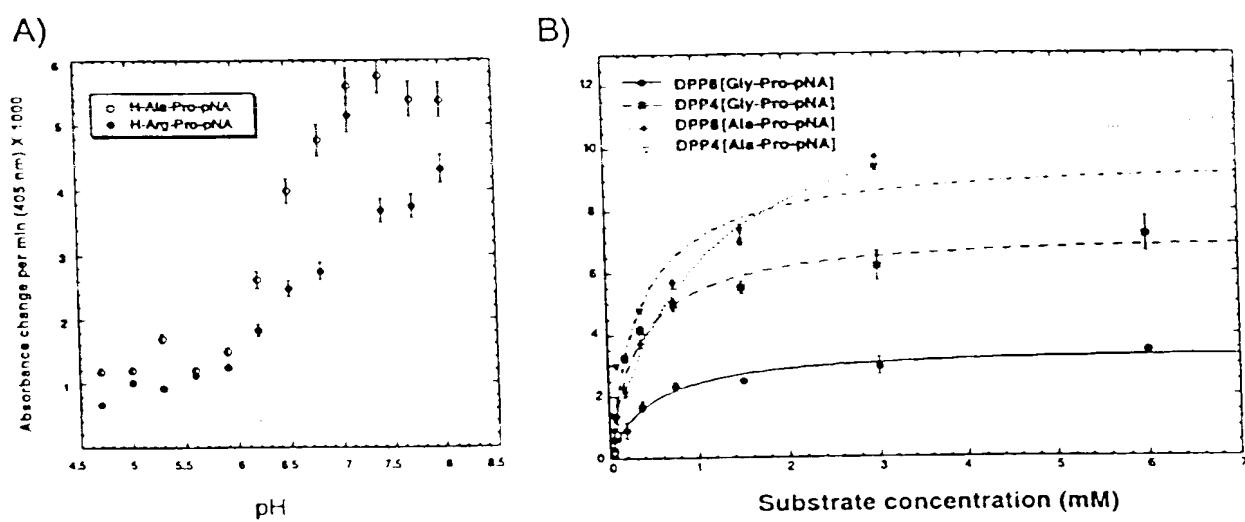


FIGURE 7

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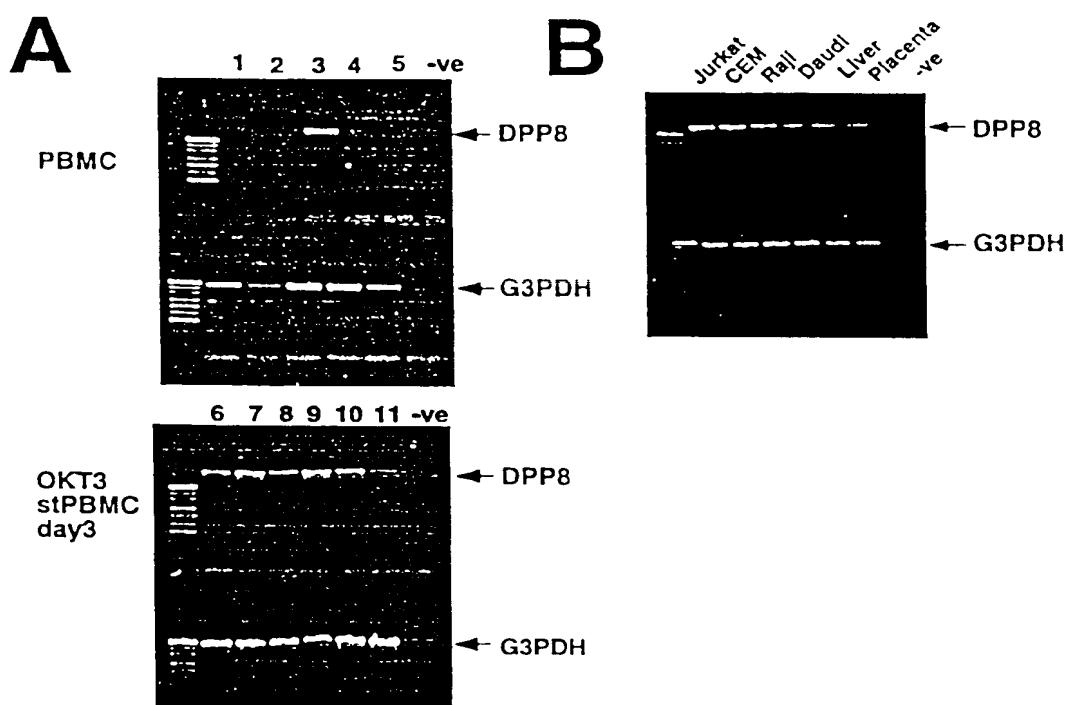


FIGURE 8

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FIGURE 9A

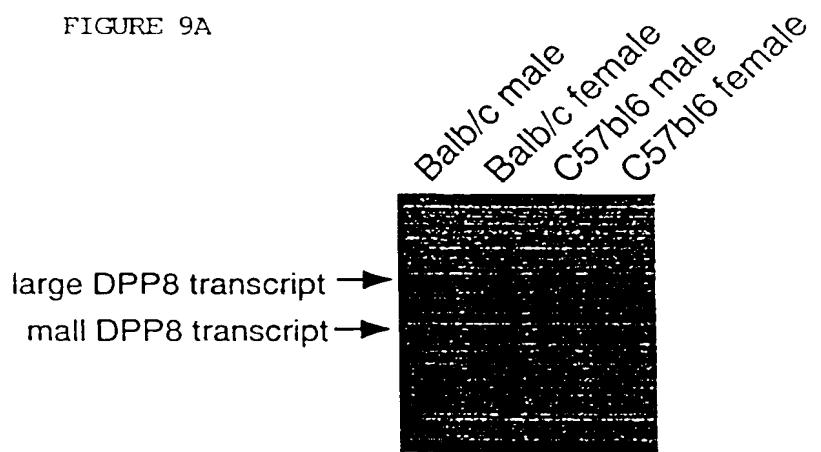


FIGURE 9B

